

EAST Search History

Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L1	6033	(sim or dreitz).inv.	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	OFF	2007/04/26 10:02
L2	1490	(t adj cell adj receptor) and canine	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	OFF	2007/04/26 10:08
L3	4	1 and 2	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	OFF	2007/04/26 10:05
L6	52749	"v.beta."\$ OR v-beta OR vb or (beta adj chain)	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	OFF	2007/04/26 10:08
L7	19530	(t adj cell adj receptor) or tcr	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	OFF	2007/04/26 10:08
L8	107588	canine or dog	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	OFF	2007/04/26 10:08
L9	512	6 7 8	US-PGPUB; USPAT; EPO; JPO; DERWENT	AND	OFF	2007/04/26 10:08

FILE 'MEDLINE' ENTERED AT 10:10:40 ON 26 APR 2007

FILE 'CAPLUS' ENTERED AT 10:10:40 ON 26 APR 2007
USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT.
PLEASE SEE "HELP USAGETERMS" FOR DETAILS.
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FILE 'BIOSIS' ENTERED AT 10:10:40 ON 26 APR 2007
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=> s sim g?/au or dreitz m?/au
L1 768 SIM G?/AU OR DREITZ M?/AU

=> s tcr or (t-cell(a)receptor)
L2 121891 TCR OR (T-CELL(A) RECEPTOR)

=> s l1 and l2
L3 68 L1 AND L2

=> s canine or dog
L4 602909 CANINE OR DOG

=> s l3 and l4
L5 7 L3 AND L4

=> dup rem l5
PROCESSING COMPLETED FOR L5
L6 4 DUP REM L5 (3 DUPLICATES REMOVED)

=> s l6 and not p/dt
MISSING TERM 'AND NOT'
The search profile that was entered contains a logical
operator followed immediately by another operator.

=> s l6 not p/dt
L7 2 L6 NOT P/DT

=> d l7 ibib ab 1-2

4-26-57

Qy	1	ATCGGACTCCTCTGTGGTGTGGCCCTTTTGTTTCCTGGGAGTAGGCCCTTTGAACGCACAA	60
Db	22	ATCAGGCTCCTCTGTCTGTGTGGCCCTTTTGTTTCCTGGCTGTAGGCCCTCGTAGATGTGAAA	81
Qy	61	GTGACTCAAACCCCGAGACAACCTCATCAAAAAAGTGGGAGCGAAAGTTTTGTTGAAATGT	120
Db	82	GTAACCCAGAGCTCGAGATATCTAGTCAAAAGGACGGGAGAGAAAGTTTTTCTGGAATGT	141
Qy	121	TCACAGAATATGGACCATGAAAGAATGTTCTGGTATCGACAAGACCCAGGTCTGGGGTTG	180
Db	142	GTCCAGGATATGGACCATGAAAAATATGTTCTGGTATCGACAAGACCCAGGTCTGGGGCTA	201
Qy	181	CGGCTGCTCTACTGGTCCTATAATATTGACAGTGTTGAGACAGGAGACATCCCTTATGGG	240
Db	202	CGGCTGATCTATTTCTCATATGATGTTAAATGAAAGAAAAAGGAGATATTCTTGAGGGG	261

Qy 241 TACAGTGTCTCGAGGAAGAAGAAGGATGCCTTCCCCTTGATTCTGGAGTCTGCTCGCATC 300
||||||| || ||||| ||| || ||||| ||| |||
Db 262 TACAGTGTCTCTAGAGAGAAGAAGGAGCGCTTCTCCCTGATTCTGGAGTCCGCCAGCACC 321

Qy 301 AACCAGACATCTGTGTACTTCTGCGCCAGCAG 332
||||||| ||||| ||| |||||
Db 322 AACCAGACATCTATGTACCTCTGTGCCAGCAG 353
<!--EndFragment-->